## APPENDIX Ia



gi3746533_human	
SEQ25_MOUSE SEQ26_Arab SEQ2_Arab SEQ14_rice SEQ16_soy SEQ22_wheat	1 MAILDSAGVTTVTENGGGEFVDLDRLRRRKSRSDSSNGLLLSGSGFNVEEDEVRLAAVS 1 MAILDSAGVTTVTENGGGEFVDLDRLRRRKSRSDSSNGLLLSGSDNNSPSDDVGAPADVR 1 MAILDSAGVTTVTENGGGEFVDLDRLRRRKSRSDSSNGLLLSGSDNNSPSDDVGAPADVR 1 MVGSDGDGDGGGGEAHAGGPRRRAGQLRGRLRDEAAPGSPPRPRPRPR 1 MAISDEPESVATALNHSSLRRRPSATSTAGLFNSPETTTDSSGDDLAKDSGSD 1 MSKGNPDPHLPGSFLPSHGGPPRKPKTPPRTFRNLPSSSTHGPAPSVAAATIAT 60
	*
gi3746533_human SEQ25_mouse	38 PDVGAAGDAPAPAPANKDGDAGVGSGHWELRCHRLQDS 41 PDLGAGGDAPAPAPAHTRDKDGRTSVGDGYWDLRCHRLQDS
SEQ2 Arab	
SEQ16_soy SEQ22_wheat	GG
	02T
gi3746533 human	WCVVMLILSNARLFLENLI
SEQ25_mouse	
SEQ26_Arab SEO2 Arab	121 PLSSDAIFKQSHAGLFNLCVVVLIAVNSRLIIENLMKYGWLIRTDFWFSSRSLRDWPL 121 PLSSDAIFKOSHAGLFNLCVVVLIAVNSRLIIENLMKYGWLIRTDFWFSSRSLRDWPL
$\overline{\text{SEQ14}}$ rice	
SEQ16_soy	97 PLSSDTIFRQSHAGLFNLCIVVLVAVNSRLIIENLMKYGWLIKSGFWFSSKSLRDWPL
SEQ22_wheat	100 PLSSDAIFRQSHAGLLNLCIVVLIAVNSRLIIENLMKYGLLIRAGFWFSARSLGDWPL

## APPENDIX Ib

* * * * * * * * * * * * * * * * * * *	194 SLLALMAHTILFLKLFSYRDVNSWCRRARAKAASAGKKASSAAAPHTVSYPDNLTYRD 203 SVFALASYSIMFLKLYSYRDVNLWCRQRRVKAKAVSTGKKVSGAAAQQAVSYPDNLTYRD 239 VTLMLLT-CIVWLKLVSYAHTSYDIRSLANAADKANPEVSYYVSLKS 239 VTLMLLT-CIVWLKLVSYAHTSYDIRSLANAADKANPEVSYYVSLKS 211 FLLIFIA-CIVWLKLVSYAHTSYDIRQLTMGGKKVDNELSTVDMDNLQPPTLGN 212 VTLMLFS-CVVWLKLVSFAHTNYDMRALTKLVEKGEALLDTLNMDYPYNVSFKS 213 FVLMFIA-SITWLKLVSFAHTNYDIRILSQSIEKGATHGSSIDEENIKGPTINS 241	** ** ** ** ** ** * * * * * * * * * *
gi3746533_human	gi3746533_human	gi3746533_human
SEQ25_mouse	SEQ25_mouse	SEQ25_mouse
SEQ26_Arab	SEQ26_Arab	SEQ26_Arab
SEQ2_Arab	SEQ2_Arab	SEQ2_Arab
SEQ14_rice	SEQ14_rice	SEQ14_rice
SEQ16_soy	SEQ16_soy	SEQ16_soy
SEQ16_soy	SEQ16_soy	SEQ16_soy

## APPENDIX IC

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	gi3746533_human SEQ25_mouse SEQ26_Arab	SEQ2_Arab SEQ14_rice	SEQ16_soy SEQ22_wheat	1			gi3746533_human	SEQ25_mouse	SEQ26_Arab	SEQ2_Arab	SEQ14_rice	SEQ16_soy	SEQ22_wheat		

## APPENDIX Id

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gi3746533_human SEQ25_mouse SEQ26_Arab SEQ2_Arab SEQ14_rice SEQ14_rice SEQ16_soy SEQ22_wheat	432 TGMP 443 TAMP 464 LGIN 443 LGIN 447 GGIN 450 SGIN 481	43 TGMMAQIPLAWFVGRFFQGNYGNAAVWLSL-IIGQPIAVLMYVHDYYVLNYEAPA 43 TAMMAQVPLAWIVGRFFQGNYGNAAVWVTL-IIGQPVAVLMYVHDYYVLNYDAPV 64 LGIMFQVPLV-FITNYLQERF-GSTVGNMIFWFIFCIFGQPMCVLLYYHDLMNRKGS 64 LGIMFQVPLV-FITNYLQERF-GSTVGNMIFWFIFCIFGQPMCVLLYYHDLMNRKGS 43 LGIMLQIPLI-VLTAYLKSKFRDTMVGNMIFWFFFCIYGQPMCVLLYYHDVMNRIEK 47 GGIMFQVPLV-LITNYLQNKFRNSMVGNMIFWFFFSILGQPMCVLLYYHDVMNRQAQ 50 SGIMFQIPLL-FLTKYLQDKFKNTMVGNMIFWFFFSIVGQPMCVLLYYHDVMNRQAQ	AV AV TVGNMIF TVGNMIF MVGNMIF	WLSL-I WVTL-I WFIFCI WFFFCI WFFFSI	IGQPIAY IGQPVAI FGQPMCY YGQPMCI LGQPMCY	7.LMYVHI 7.LMYVHI 7.LLY	DYYVLNYEAPA DYYVLNYDAPV -YHDLMNRKGS -YHDLMNRKGS -YHDVMNRIEK -YHDLMNRKGK -YHDVMNRIEK -YHDVMNRQAQ
gi3746533_human SEQ25_mouse SEQ26_Arab SEQ2_Arab SEQ14_rice SEQ16_soy SEQ22_wheat	486 AEA 497 G-V 519 MS- 519 AS- 499 AR- 503 LD- 506 TNG	488 498 520 500 504 508					